



amdt a

SEQUENCE LISTING

<110> Wood, Clive
Chaudhary, Divya
Long, Andrew
Genetics Institute, Inc.

<120> TRADE MOLECULES, AND USES RELATED THERETO

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 Glu Ser Gly Ala Ile Ile His Pro Ala Thr Gln Thr Ser Leu Gln Glu
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ggc tat ggg gag gat gca cag tgt gtg acg tgc cgg ctg cac agg ttc 240
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Glu Ser Gly Ala Ile Ile His Pro Ala Thr Gln Thr Ser Leu Gln Val			
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 Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
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 Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Leu Asp Arg Pro
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 Gln Leu His Glu Tyr Ala His Arg Ala Cys Cys Gln Cys Arg Arg Asp
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 Val Pro Thr Phe Phe Gly Ser Leu Thr Gln Ser Ile Cys Gly Glu Phe
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 Ser Phe Cys Asp Ser Tyr Pro Glu Leu Ala Gly Glu Asp Ile His Ser
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Gly Tyr Gly Glu Asp Ala Gln Cys Val Pro Cys Arg Pro His Arg Phe	
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Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Ala Asp Cys Ala	
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Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro	
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195 200 205	

cag gac att cag tac aat ggc tct gag ctg tca tgc ttt gac cag cct	672
Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Phe Asp Gln Pro	
210 215 220	
cgg ctc cgc cac tgt gcc cat aga gca tgc tgt cag tat cac cgg gac	720
Arg Leu Arg His Cys Ala His Arg Ala Cys Cys Gln Tyr His Arg Asp	
225 230 235 240	
tca gcc cca atg tat ggg cct gtt cac ctg att ccg tcc ttg tgc tgt	768
Ser Ala Pro Met Tyr Gly Pro Val His Leu Ile Pro Ser Leu Cys Cys	
245 250 255	
gaa gag gcc cgc agc tct gcc cga gct gtg ctt ggc tgt ggg ctg cgt	816
Glu Glu Ala Arg Ser Ser Ala Arg Ala Val Leu Gly Cys Gly Leu Arg	
260 265 270	
tct ccc act acc ctc cag gag aga aac ccg gct tct gtg ggg gac acg	864
Ser Pro Thr Thr Leu Gln Glu Arg Asn Pro Ala Ser Val Gly Asp Thr	
275 280 285	
atg cca gcc ttc ttc ggg tct gtt tcc cgt tcc atc tgc gct gaa ttt	912
Met Pro Ala Phe Phe Gly Ser Val Ser Arg Ser Ile Cys Ala Glu Phe	
290 295 300	
tct gat gcc tgg cct ctg atg cag aat cct ctg ggt ggt gac agc tct	960
Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Leu Gly Gly Asp Ser Ser	
305 310 315 320	
ctc tgt gac tct tat cct gaa ctc act gga gaa gat acc aat tcc ctc	1008
Leu Cys Asp Ser Tyr Pro Glu Leu Thr Gly Glu Asp Thr Asn Ser Leu	
325 330 335	
aat ccc gaa aac gaa agc gca gca tct ctg gat tcc agt ggc ggc cag	1056
Asn Pro Glu Asn Glu Ser Ala Ala Ser Leu Asp Ser Ser Gly Gly Gln	
340 345 350	
gat ctg gct ggg aca gct gct cta gag tct tct ggg aat gtt tca gaa	1104
Asp Leu Ala Gly Thr Ala Ala Leu Glu Ser Ser Gly Asn Val Ser Glu	
355 360 365	
tct act gac tca cct aga cat ggt gac act ggt aca gtc tgg gag cag	1152
Ser Thr Asp Ser Pro Arg His Gly Asp Thr Gly Thr Val Trp Glu Gln	
370 375 380	
acg cta gct cag gat gct caa agg act cca agc caa gga ggc tgg gaa	1200
Thr Leu Ala Gln Asp Ala Gln Arg Thr Pro Ser Gln Gly Gly Trp Glu	
385 390 395 400	
gac agg gaa aac ctg aat cta gcc atg ccc aca gcc ttc cag gat gcc	1248
Asp Arg Glu Asn Leu Asn Leu Ala Met Pro Thr Ala Phe Gln Asp Ala	
405 410 415	
tgaaggccat cttcctgacg tggaggtgtg ggtctggaca agcctgtgat gaggcctaca	1308
gactgagcag tcttggtgtc tggaagcaaa aataaatctg aaccaaactg acaacatttc	1368
catcctttca gccactagct tctgagccag accagctgta agctgaaacc ccagcaagaa	1428

gcaaggagag actgactgta ggcggccttg ggacatgtgc ttcttccta agcgagaacc 1488
 ttagctgggg ccaatttgaa ggacccatgg gtggaatgtg ctgcctgtga gcttggtggc 1548
 acagcaggac ccagcctggc tccttcttat gtccacgggtg aatgtggttt cacaagaccc 1608
 agagtataaa ctttcataga cattctcttt tagaaataat ccattaccct gtcttcaaaa 1668
 accaaaaaaaa aaaaaaagtg gtgttaaggt tttgaacatc acctagccaa gttagtaaaa 1728
 tctttatttg tatttcatct caattttttt aactattcat tttccttgta tgaattcttg 1788
 tgtgttttat gtgtaaatat attcattatt ttgacactat caatattctt tgtgggtttg 1848
 taatttttac ttttattaat gactcaagct gtaaaaataa actaatttca acgtcgacgc 1908
 ggccgc 1914

<210> 6
 <211> 416
 <212> PRT
 <213> Mus musculus

<400> 6
 Met Ala Leu Lys Val Leu Pro Leu His Arg Thr Val Leu Phe Ala Ala
 1 5 10 15
 Ile Leu Phe Leu Leu His Leu Ala Cys Lys Val Ser Cys Glu Thr Gly
 20 25 30
 Asp Cys Arg Gln Gln Glu Phe Lys Asp Arg Ser Gly Asn Cys Val Leu
 35 40 45
 Cys Lys Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
 50 55 60
 Gly Tyr Gly Glu Asp Ala Gln Cys Val Pro Cys Arg Pro His Arg Phe
 65 70 75 80
 Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Ala Asp Cys Ala
 85 90 95
 Leu Val Asn Arg Phe Gln Arg Ala Asn Cys Ser His Thr Ser Asp Ala
 100 105 110
 Val Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
 115 120 125
 Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
 130 135 140
 Tyr Glu Pro His Cys Thr Ser Lys Val Asn Leu Val Lys Ile Ser Ser
 145 150 155 160
 Thr Val Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser

165					170					175					
Ala	Leu	Ala	Thr	Val	Leu	Leu	Ala	Leu	Leu	Ile	Leu	Cys	Val	Ile	Tyr
			180					185					190		
Cys	Lys	Arg	Gln	Phe	Met	Glu	Lys	Lys	Pro	Ser	Trp	Ser	Leu	Arg	Ser
		195					200					205			
Gln	Asp	Ile	Gln	Tyr	Asn	Gly	Ser	Glu	Leu	Ser	Cys	Phe	Asp	Gln	Pro
	210					215					220				
Arg	Leu	Arg	His	Cys	Ala	His	Arg	Ala	Cys	Cys	Gln	Tyr	His	Arg	Asp
225					230					235					240
Ser	Ala	Pro	Met	Tyr	Gly	Pro	Val	His	Leu	Ile	Pro	Ser	Leu	Cys	Cys
			245						250					255	
Glu	Glu	Ala	Arg	Ser	Ser	Ala	Arg	Ala	Val	Leu	Gly	Cys	Gly	Leu	Arg
		260						265					270		
Ser	Pro	Thr	Thr	Leu	Gln	Glu	Arg	Asn	Pro	Ala	Ser	Val	Gly	Asp	Thr
		275					280					285			
Met	Pro	Ala	Phe	Phe	Gly	Ser	Val	Ser	Arg	Ser	Ile	Cys	Ala	Glu	Phe
	290					295					300				
Ser	Asp	Ala	Trp	Pro	Leu	Met	Gln	Asn	Pro	Leu	Gly	Gly	Asp	Ser	Ser
305					310					315					320
Leu	Cys	Asp	Ser	Tyr	Pro	Glu	Leu	Thr	Gly	Glu	Asp	Thr	Asn	Ser	Leu
		325							330				335		
Asn	Pro	Glu	Asn	Glu	Ser	Ala	Ala	Ser	Leu	Asp	Ser	Ser	Gly	Gly	Gln
		340						345					350		
Asp	Leu	Ala	Gly	Thr	Ala	Ala	Leu	Glu	Ser	Ser	Gly	Asn	Val	Ser	Glu
	355						360				365				
Ser	Thr	Asp	Ser	Pro	Arg	His	Gly	Asp	Thr	Gly	Thr	Val	Trp	Glu	Gln
	370					375					380				
Thr	Leu	Ala	Gln	Asp	Ala	Gln	Arg	Thr	Pro	Ser	Gln	Gly	Gly	Trp	Glu
385					390					395					400
Asp	Arg	Glu	Asn	Leu	Asn	Leu	Ala	Met	Pro	Thr	Ala	Phe	Gln	Asp	Ala
			405						410				415		

<210> 7

<211> 27

<212> DNA

<213> Mus musculus

<400> 7

aggccatctt cctgacgtgg aggtgtg

27

<210> 8
 <211> 35
 <212> DNA
 <213> Mus musculus

<400> 8
 cggaattcgt ttcagctcag cacattccaa ggccg

35

<210> 9
 <211> 9
 <212> PRT
 <213> Homo sapiens

<400> 9
 Ser Thr Ala Ser Ser Pro Arg Asp Thr
 1 5

<210> 10
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 10
 Asp Lys Thr His Thr Cys Pro
 1 5

<210> 11
 <211> 417
 <212> PRT
 <213> Homo sapiens

<400> 11
 Met Ala Leu Lys Val Leu Leu Glu Gln Glu Lys Thr Phe Phe Thr Leu
 1 5 10 15

Leu Val Leu Leu Gly Tyr Leu Ser Cys Lys Val Thr Cys Glu Ser Gly
 20 25 30

Asp Cys Arg Gln Gln Glu Phe Arg Asp Arg Ser Gly Asn Cys Val Pro
 35 40 45

Cys Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
 50 55 60

Gly Tyr Gly Glu Asp Ala Gln Cys Val Thr Cys Arg Leu His Arg Phe
 65 70 75 80

Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Leu Asp Cys Ala
 85 90 95

Val Val Asn Arg Phe Gln Lys Ala Asn Cys Ser Ala Thr Ser Asp Ala
 100 105 110

Ile Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
 115 120 125

Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
 130 135 140
 Tyr Glu Pro His Cys Ala Ser Lys Val Asn Leu Val Lys Ile Ala Ser
 145 150 155 160
 Thr Ala Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
 165 170 175
 Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
 180 185 190
 Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
 195 200 205
 Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Phe Asp Arg Pro
 210 215 220
 Gln Leu His Glu Tyr Ala His Arg Ala Cys Cys Gln Cys Arg Arg Asp
 225 230 235 240
 Ser Val Gln Thr Cys Gly Pro Val Arg Leu Leu Pro Ser Met Cys Cys
 245 250 255
 Glu Glu Ala Cys Ser Pro Asn Pro Ala Thr Leu Gly Cys Gly Val His
 260 265 270
 Ser Ala Ala Ser Leu Gln Ala Arg Asn Ala Gly Pro Ala Gly Glu Met
 275 280 285
 Val Pro Thr Phe Phe Gly Ser Leu Thr Gln Ser Ile Cys Gly Glu Phe
 290 295 300
 Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Met Gly Gly Asp Asn Ile
 305 310 315 320
 Ser Phe Cys Asp Ser Tyr Pro Glu Leu Thr Gly Glu Asp Ile His Ser
 325 330 335
 Leu Asn Pro Glu Leu Glu Ser Ser Thr Ser Leu Asp Ser Asn Ser Ser
 340 345 350
 Gln Asp Leu Val Gly Gly Ala Val Pro Val Gln Ser His Ser Glu Asn
 355 360 365
 Phe Thr Ala Ala Thr Asp Leu Ser Arg Tyr Asn Asn Thr Leu Val Glu
 370 375 380
 Ser Ala Ser Thr Gln Asp Ala Leu Thr Met Arg Ser Gln Leu Asp Gln
 385 390 395 400
 Glu Ser Gly Ala Val Ile His Pro Ala Thr Gln Thr Ser Leu Gln Glu
 405 410 415
 Ala

<210> 12
 <211> 423
 <212> PRT
 <213> Homo sapiens

<400> 12
 Met Ala Leu Lys Val Leu Leu Glu Gln Glu Lys Thr Phe Phe Thr Leu
 1 5 10 15
 Leu Val Leu Leu Gly Tyr Leu Ser Cys Lys Val Thr Cys Glu Ser Gly
 20 25 30
 Asp Cys Arg Gln Gln Glu Phe Arg Asp Arg Ser Gly Asn Cys Val Pro
 35 40 45
 Cys Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
 50 55 60
 Gly Tyr Gly Glu Asp Ala Gln Cys Val Ala Cys Arg Leu His Arg Phe
 65 70 75 80
 Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Leu Asp Cys Ala
 85 90 95
 Val Val Asn Arg Phe Gln Lys Ala Asn Cys Ser Ala Thr Ser Asp Ala
 100 105 110
 Ile Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
 115 120 125
 Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
 130 135 140
 Tyr Glu Pro His Cys Ala Ser Lys Val Asn Leu Val Lys Ile Ala Ser
 145 150 155 160
 Thr Ala Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
 165 170 175
 Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
 180 185 190
 Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
 195 200 205
 Gln Asp Ile Gln Tyr Asn Glu Ser Glu Leu Ser Cys Phe Asp Arg Pro
 210 215 220
 Gln Leu His Glu Tyr Ala His Arg Ala Cys Cys Gln Cys Arg Arg Asp
 225 230 235 240
 Ser Val Gln Thr Cys Gly Pro Val Arg Leu Leu Pro Ser Met Cys Cys
 245 250 255
 Glu Glu Ala Cys Ser Pro Asn Pro Ala Thr Leu Gly Cys Gly Val His

260	265	270
Ser Ala Ala Ser Leu Gln Ala Arg Asn Ala Gly Pro Ala Gly Glu Met		
275	280	285
Val Pro Thr Phe Phe Gly Ser Leu Thr Gln Ser Ile Cys Gly Glu Phe		
290	295	300
Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Met Gly Gly Asp Asn Ile		
305	310	315
Ser Phe Cys Asp Ser Tyr Pro Glu Leu Thr Gly Glu Asp Ile His Ser		
325	330	335
Leu Asn Pro Glu Leu Glu Ser Ser Thr Ser Leu Asp Ser Asn Ser Ser		
340	345	350
Gln Asp Leu Val Gly Gly Ala Val Pro Val Gln Ser His Ser Glu Asn		
355	360	365
Phe Thr Ala Ala Thr Asp Leu Ser Arg Tyr Asn Asn Thr Leu Val Glu		
370	375	380
Ser Ala Ser Thr Gln Asp Ala Leu Thr Met Arg Ser Gln Leu Asp Gln		
385	390	395
Glu Ser Gly Ala Val Ile His Pro Ala Thr Gln Thr Ser Leu Gln Glu		
405	410	415
Arg Gln Arg Leu Gly Ser Leu		
420		

<210> 13
 <211> 420
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:Consensus
 sequence

<400> 13
Met Ala Leu Lys Val Leu Leu Glu Gln Glu Lys Thr Phe Phe Thr Leu
1 5 10 15
Leu Val Leu Leu Gly Tyr Leu Ser Cys Lys Val Thr Cys Glu Ser Gly
20 25 30
Asp Cys Arg Gln Gln Glu Phe Arg Asp Arg Ser Gly Asn Cys Val Pro
35 40 45
Cys Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
50 55 60
Gly Tyr Gly Glu Asp Ala Gln Cys Val Thr Cys Arg Leu His Arg Phe
65 70 75 80

Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Leu Asp Cys Ala
 85 90 95

Val Val Asn Arg Phe Gln Lys Ala Asn Cys Ser Ala Thr Ser Asp Ala
 100 105 110

Ile Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
 115 120 125

Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
 130 135 140

Tyr Glu Pro His Cys Ala Ser Lys Val Asn Leu Val Lys Ile Ala Ser
 145 150 155 160

Thr Ala Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
 165 170 175

Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
 180 185 190

Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
 195 200 205

Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Phe Asp Arg Pro
 210 215 220

Gln Leu His Glu Tyr Ala His Arg Ala Cys Cys Gln Cys Arg Arg Asp
 225 230 235 240

Ser Val Gln Thr Cys Gly Pro Val Arg Leu Leu Pro Ser Met Cys Cys
 245 250 255

Glu Glu Ala Cys Ser Pro Asn Pro Ala Thr Leu Gly Cys Gly Val His
 260 265 270

Ser Ala Ala Ser Leu Gln Ala Arg Asn Ala Gly Pro Ala Gly Glu Met
 275 280 285

Val Pro Thr Phe Phe Gly Ser Leu Thr Gln Ser Ile Cys Gly Glu Phe
 290 295 300

Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Met Gly Gly Asp Asn Ile
 305 310 315 320

Ser Phe Cys Asp Ser Tyr Pro Glu Leu Thr Gly Glu Asp Ile His Ser
 325 330 335

Leu Asn Pro Glu Leu Glu Ser Ser Thr Ser Leu Asp Ser Asn Ser Ser
 340 345 350

Gln Asp Leu Val Gly Gly Ala Val Pro Val Gln Ser His Ser Glu Asn
 355 360 365

Phe Thr Ala Ala Thr Asp Leu Ser Arg Tyr Asn Asn Thr Leu Val Glu
 370 375 380

Ser Ala Ser Thr Gln Asp Ala Leu Thr Met Arg Ser Gln Leu Asp Gln
 385 390 395 400

Glu Ser Gly Ala Ile His Pro Ala Thr Gln Thr Ser Leu Gln Gln Arg
 405 410 415

Leu Gly Ser Leu
 420

<210> 14
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 14
 Glu Ser Gly Ala Ile Ile His Pro Ala Thr Gln Thr Ser Leu Gln Glu
 1 5 10 15

Ala

<210> 15
 <211> 23
 <212> PRT
 <213> Homo sapiens

<400> 15
 Glu Ser Gly Ala Ile Ile His Pro Ala Thr Gln Thr Ser Leu Gln Val
 1 5 10 15

Arg Gln Arg Leu Gly Ser Leu
 20

<210> 16
 <211> 34
 <212> PRT
 <213> Homo sapiens

<400> 16
 Cys Pro Thr Gly Leu Tyr Thr His Ser Gly Glu Cys Cys Lys Ala Cys
 1 5 10 15

Asn Leu Gly Glu Gly Val Ala Gln Pro Cys Gly Ala Asn Gln Thr Val
 20 25 30

Cys Glu

<210> 17
 <211> 34
 <212> PRT
 <213> Homo sapiens

<400> 17
 Cys Gly Asp Thr Tyr Pro Ser Asn Asp Arg Cys Cys His Glu Cys Arg
 1 5 10 15
 Pro Gly Asn Gly Met Val Ser Arg Cys Ser Arg Ser Gln Asn Thr Val
 20 25 30
 Cys Arg

<210> 18
 <211> 35
 <212> PRT
 <213> Homo sapiens

<400> 18
 Cys Arg Glu Lys Gln Tyr Leu Leu Asn Ser Gln Cys Cys Ser Leu Cys
 1 5 10 15
 Gln Pro Gly Gln Lys Leu Val Ser Asp Cys Thr Glu Phe Thr Glu Thr
 20 25 30
 Glu Cys Leu
 35

<210> 19
 <211> 40
 <212> PRT
 <213> Homo sapiens

<400> 19
 Cys Arg Gln Gln Glu Phe Arg Asp Arg Ser Gly Asn Cys Val Pro Cys
 1 5 10 15
 Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe Gly
 20 25 30
 Tyr Gly Glu Asp Ala Gln Cys Val
 35 40

<210> 20
 <211> 42
 <212> PRT
 <213> Homo sapiens

<400> 20
 Pro Cys Leu Asp Ser Val Thr Phe Ser Asp Val Val Ser Ala Thr Glu
 1 5 10 15
 Pro Cys Lys Pro Cys Thr Glu Cys Val Gly Leu Gln Ser Met Ser Ala
 20 25 30
 Pro Cys Val Glu Ala Asp Asp Ala Val Cys

35

40

<210> 21
 <211> 42
 <212> PRT
 <213> Homo sapiens

<400> 21
 Pro Cys Gly Pro Gly Phe Tyr Asn Asp Val Val Ser Ser Lys Pro Cys
 1 5 10 15
 Lys Pro Cys Thr Trp Cys Asn Leu Arg Ser Gly Ser Glu Arg Lys Gln
 20 25 30
 Leu Cys Thr Ala Thr Gln Asp Thr Val Cys
 35 40

<210> 22
 <211> 43
 <212> PRT
 <213> Homo sapiens

<400> 22
 Pro Cys Gly Glu Ser Glu Phe Leu Asp Thr Trp Asn Arg Glu Thr His
 1 5 10 15
 Cys His Gln His Lys Tyr Cys Asp Pro Asn Leu Gly Leu Arg Val Gln
 20 25 30
 Gln Lys Gly Thr Ser Glu Thr Asp Thr Ile Cys
 35 40

<210> 23
 <211> 41
 <212> PRT
 <213> Homo sapiens

<400> 23
 Thr Cys Arg Leu His Arg Phe Lys Glu Asp Trp Gly Phe Gln Lys Cys
 1 5 10 15
 Lys Pro Cys Leu Asp Cys Ala Val Val Asn Arg Phe Gln Lys Ala Asn
 20 25 30
 Cys Ser Ala Thr Ser Asp Ala Ile Cys
 35 40

<210> 24
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 24

Asp Lys Thr His Thr Cys Pro
1 5

Al
concord